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RESULT 4
QUERY Match 94.9% Score 1972 Length 144
Host Local Similarity 94.9% Prod. No. 42614
Matches 282 Conserved 19 Mismatches 19 Indels 0 Gaps 0

1 CAVRTIGANKHIVATGNNVYVHIVHNSV 40
II 115 CAVRTIGANKHIVATGNNVYVHIVHNSV 144
FT NON_TER 1 19 POTENTIAL
STRAIN 20 142 PANCREATIC RIBONUCLEASE
SEQUENCE 144 AA: 15755 MW: 11190400.3470E CR644

QUERY Match 94.9% Score 1972 Length 144
Host Local Similarity 94.9% Prod. No. 42614
Matches 282 Conserved 19 Mismatches 19 Indels 0 Gaps 0

1 CAVRTIGANKHIVATGNNVYVHIVHNSV 40
II 115 CAVRTIGANKHIVATGNNVYVHIVHNSV 144
FT NON_TER 1 19 POTENTIAL
STRAIN 20 142 PANCREATIC RIBONUCLEASE
SEQUENCE 144 AA: 15755 MW: 11190400.3470E CR644

QUERY Match 94.9% Score 1972 Length 144
Host Local Similarity 94.9% Prod. No. 42614
Matches 282 Conserved 19 Mismatches 19 Indels 0 Gaps 0

1 CAVRTIGANKHIVATGNNVYVHIVHNSV 40
II 115 CAVRTIGANKHIVATGNNVYVHIVHNSV 144
FT NON_TER 1 19 POTENTIAL
STRAIN 20 142 PANCREATIC RIBONUCLEASE
SEQUENCE 144 AA: 15755 MW: 11190400.3470E CR644

QUERY Match 94.9% Score 1972 Length 144
Host Local Similarity 94.9% Prod. No. 42614
Matches 282 Conserved 19 Mismatches 19 Indels 0 Gaps 0

1 CAVRTIGANKHIVATGNNVYVHIVHNSV 40
II 115 CAVRTIGANKHIVATGNNVYVHIVHNSV 144
FT NON_TER 1 19 POTENTIAL
STRAIN 20 142 PANCREATIC RIBONUCLEASE
SEQUENCE 144 AA: 15755 MW: 11190400.3470E CR644

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RESULT 5
QUERY Match 84.9% Score 141 Length 164
Host Local Similarity 84.9% Prod. No. 42614
Matches 252 Conserved 19 Mismatches 32 Indels 0 Gaps 0

1 CAVRTIGANKHIVATGNNVYVHIVHNSV 40
II 115 CAVRTIGANKHIVATGNNVYVHIVHNSV 145
FT NON_TER 1 19 POTENTIAL
STRAIN 20 142 PANCREATIC RIBONUCLEASE
SEQUENCE 145 AA: 18997 MW: 83167654A982A1CF CR644

QUERY Match 84.9% Score 141 Length 164
Host Local Similarity 84.9% Prod. No. 42614
Matches 252 Conserved 19 Mismatches 32 Indels 0 Gaps 0

1 CAVRTIGANKHIVATGNNVYVHIVHNSV 40
II 115 CAVRTIGANKHIVATGNNVYVHIVHNSV 145
FT NON_TER 1 19 POTENTIAL
STRAIN 20 142 PANCREATIC RIBONUCLEASE
SEQUENCE 145 AA: 18997 MW: 83167654A982A1CF CR644

QUERY Match 84.9% Score 141 Length 164
Host Local Similarity 84.9% Prod. No. 42614
Matches 252 Conserved 19 Mismatches 32 Indels 0 Gaps 0

1 CAVRTIGANKHIVATGNNVYVHIVHNSV 40
II 115 CAVRTIGANKHIVATGNNVYVHIVHNSV 145
FT NON_TER 1 19 POTENTIAL
STRAIN 20 142 PANCREATIC RIBONUCLEASE
SEQUENCE 145 AA: 18997 MW: 83167654A982A1CF CR644

QUERY Match 84.9% Score 141 Length 164
Host Local Similarity 84.9% Prod. No. 42614
Matches 252 Conserved 19 Mismatches 32 Indels 0 Gaps 0

1 CAVRTIGANKHIVATGNNVYVHIVHNSV 40
II 115 CAVRTIGANKHIVATGNNVYVHIVHNSV 145
FT NON_TER 1 19 POTENTIAL
STRAIN 20 142 PANCREATIC RIBONUCLEASE
SEQUENCE 145 AA: 18997 MW: 83167654A982A1CF CR644

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